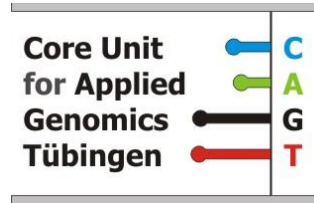


Sample Sheet



Next-Generation-Sequencing

Important parameters for successful sequencing

- **Before you will send the samples, please contact us (07071 29 86297; Email: peter.bauer@med.uni-tuebingen.de).**
- We can't start with the project, when we don't have the sample sheet.
- A minimum of 3 µg genomic DNA (> 100 ng/µL) is required. For enrichment projects, there can be other specifications. We also accept **fresh** EDTA-blood samples or cultured cells (but should know shipment dates in advance). If you cannot provide 3 µg gDNA please contact us immediately.
- After finishing the project we can send the samples back to you, at your expense.
- We will store raw data files for a maximum of 12 months, but you can get a copy of your dataset when providing a HDD drive.
- DNA stocks and sequencing libraries will be stored for 12 months and discarded afterwards without further notice

Contact us:

Core Unit for Applied Genomics

Medical Genetics

Calwerstr. 7

D 72076 Tübingen

Phone: 07071 29 77692 (head) / 07071 29 86297 (lab)

Email: peter.bauer@med.uni-tuebingen.de

Sample Sheet



Contact information

Institut		Name	
Department		Head	
Street / Number		Email	
ZIP, City		Phone	
Country		Fax	

General Sample information

Species	Source (e.g. blood, brain ..)
<input type="checkbox"/> DNA <input type="checkbox"/> RNA <input type="checkbox"/> cDNA <input type="checkbox"/> native <input type="checkbox"/> Average frag size < 1000bp	Prep. method: Elution buffer: OD measurd by: Nanodrop OD260/280 >1,8; OD260/230 > 2,0
<input type="checkbox"/> Amplicons	Cleanup kit: Elution buffer:
<input type="checkbox"/> Agarose gel appended	

Internal use

Job number	
Project ID	
Date / localization	
Quote ID	

Sample Sheet

Core Unit for Applied Genomics Tübingen

- C
- A
- G
- T

No	Sample name	Conc. (ng/ μ L)	OD 260/280	OD 260/230	Total vol (μ L)	Total amount (μ g)	Comment
1							
2							
3							
4							
5							
6							
7							
8							
9							
10							